

Resource Summary Report

Generated by [kravitz2](#) on Aug 2, 2026

ZENODO

RRID:SCR_004129

Type: Tool

Proper Citation

ZENODO (RRID:SCR_004129)

Resource Information

URL: <https://zenodo.org/>

Proper Citation: ZENODO (RRID:SCR_004129)

Description: Repository for all research outputs from across all fields of science in any file format as well as both positive and negative results. They assign all publicly available uploads a Digital Object Identifier (DOI) to make the upload easily and uniquely citeable. They further support harvesting of all content via the OAI-PMH protocol. They promote peer-reviewed openly accessible research, and curate uploads. ZENODO allows users to create their own collection and accept or reject all uploads to it. They allow for uploading under a multitude of different licenses and access levels.

Abbreviations: ZENODO

Synonyms: Zenodo

Resource Type: catalog, software repository, software resource, data or information resource, database, data repository, service resource, storage service resource

Keywords: data set, software resource, video resource, audio track, image, poster, presentation, publication, digital preservation, digital archive, persistent identifier, digital object identifier, openaire orphan record repository

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Resource Name: ZENODO

Resource ID: SCR_004129

Alternate IDs: DOI:10.5281, nlx_158614, DOI:10.17616/R3QP53, DOI:10.25504/FAIRsharing.wy4egf, r3d100011858

Alternate URLs: <https://doi.org/10.17616/R3QP53>, <https://doi.org/10.17616/r3QP53>, <https://doi.org/10.5281/>, <https://dx.doi.org/10.5281/>,

<https://fairsharing.org/10.25504/FAIRsharing.wy4egf>, <https://doi.org/10.17616/R35W56>

Record Creation Time: 20220129T080222+0000

Record Last Update: 20260803T040353+0000

Ratings and Alerts

No rating or validation information has been found for ZENODO.

No alerts have been found for ZENODO.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 4900 mentions in open access literature.

Listed below are recent publications. The full list is available at [kravitz2](#).

Bettinelli A, et al. (2026) Assessing the agreement of radiomic tools for dosiomics: A multi-software comparative study. Medical physics, 53(1), e70203.

Daniel MM, et al. (2026) Integrative Bioinformatics Reveals Novel Molecular Mechanisms and Therapeutic Targets in Acute Myeloid Leukaemia. Journal of cellular and molecular medicine, 30(1), e71007.

Rotem A, et al. (2026) Differentiation between regulated and disrupted growth arrests allows tailoring of effective treatments for antibiotic persistence. Science advances, 12(1), eadt6577.

Kot E, et al. (2026) Crystal structure of fatty acid thioesterase A bound by 129 fragments provides diverse development opportunities. Pest management science, 82(1), 151.

Martin-Gonzalez E, et al. (2026) Multi-Ancestry Epigenome-Wide Association Study of Asthma Exacerbations. Allergy, 81(1), 263.

Berselli A, et al. (2026) The effect of ionic strength on PETase enzymes: An experimental and computational study. Protein science : a publication of the Protein Society, 35(1), e70386.

Li X, et al. (2026) Talin-tensin3 interactions regulate fibrillar adhesion formation and tensin3 phase separation. The Journal of cell biology, 225(1).

Zanini A, et al. (2026) Mapping functional homologies between human and marmoset brain networks using movie-driven ultra-high field fMRI. Communications biology, 9(1), 21.

Saviluoto A, et al. (2026) Association of Cerebral Oxygenation During Prehospital Anaesthesia and Functional Outcome: A Prospective, Observational Multi-Centre Cohort Study of 1014 Patients. *Acta anaesthesiologica Scandinavica*, 70(1), e70161.

Andonie CR, et al. (2026) Comparative Analysis of Glucagon Receptor Agonists vs. Resmetirom in MASLD and MASH: Network Meta-Analysis of Clinical Trials. *Endocrinology, diabetes & metabolism*, 9(1), e70157.

Yilmaz V, et al. (2026) Comparative Transcriptomic Profiling Identifies IL-1 β as a Non-Canonical Driver of Human Dendritic Cell Maturation. *Iranian journal of biotechnology*, 24(1), e4185.

Zhao Z, et al. (2026) Multi-omics analysis unveils tumor heterogeneity and immunotherapy predictive model in breast cancer for precision medicine and early detection. *Neoplasia* (New York, N.Y.), 71, 101260.

Yan Y, et al. (2026) A novel mouse cell line model reveals the tumor intrinsic and immune characteristics of EGFR-mutant lung cancer. *Theranostics*, 16(2), 1006.

Zappon E, et al. (2026) An efficient end-to-end computational framework for the generation of ECG calibrated volumetric models of human atrial electrophysiology. *Medical image analysis*, 107(Pt B), 103822.

Freudenthal J, et al. (2026) A Novel Protistan Trait Database Reveals Functional Redundancy and Complementarity in Terrestrial Protists (Amoebozoa and Rhizaria). *Molecular ecology resources*, 26(1), e70064.

Verna G, et al. (2026) A missense mutation in Muc2 promotes gut microbiome and metabolome-dependent colitis-associated tumorigenesis. *The Journal of clinical investigation*, 136(1).

Widomski C, et al. (2026) Beyond the emm-type: fine-tuning Group A Streptococcus typing with Enn and Mrp. *Microbiology spectrum*, 14(1), e0204725.

Michel M, et al. (2025) Noninvasive Multicancer Detection Using DNA Hypomethylation of LINE-1 Retrotransposons. *Clinical cancer research : an official journal of the American Association for Cancer Research*, 31(7), 1275.

Roca MS, et al. (2025) Valproic acid improves the efficacy of oxaliplatin/fluoropyrimidine-based chemotherapy by targeting cancer stem cell via β -Catenin modulation in colorectal cancer. *Cell death & disease*, 16(1), 583.

Kondi $\ddot{S}$ T, et al. (2025) Shinyscreen: mass spectrometry data inspection and quality checking utility. *Journal of cheminformatics*, 17(1), 98.